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TC 1700
SEQ ID NO: 11

	<u>Peptide</u>	<u>Sequence</u>	
P1 ₍₂₈₀₋₂₉₃₎	AlaLeuAspThrAsnTyrCysPheSerSerThrGluLysAsn		
P2 ₍₂₈₄₋₂₉₇₎	AsnTyrCysSerSerThrGluLysAsnCysCysValArg		SEQ ID NO: 12
P3 ₍₂₈₈₋₃₀₁₎	SerSerThrGluLysAsnCysCysValArgGlnLeuTyrIle		SEQ ID NO: 13
P4 ₍₂₉₄₋₃₀₇₎	CysCysValArgGlnLeuTyrIleAspPheArgLysAspLeu		SEQ ID NO: 14
P5 ₍₂₉₈₋₃₁₁₎	GlnLeuTyrIleAspPheArgLysAspLeuGlyTrpLysTrp		SEQ ID NO: 15
P6 ₍₃₀₂₋₃₁₅₎	AspPheArgLysAspLeuGlyTrpLysTrpIleHisGluPro		SEQ ID NO: 16
P7 ₍₃₀₆₋₃₁₉₎	AspLeuGlyTrpLysTrpIleHisGluProLysGlyTyrHis		SEQ ID NO: 17
P8 ₍₃₀₈₋₃₂₁₎	GlyTrpLysTrpIleHisGluProLysGlyTyrHisAlaAsn		SEQ ID NO: 18
P9 ₍₃₁₂₋₃₂₅₎	IleHisGluProLysGlyTyrHisAlaAsnPheCysLeuGly		SEQ ID NO: 19
P10 ₍₃₁₆₋₃₂₉₎	LysGlyTyrHisAlaAsnPheCysLeuGlyProCysProTyr		SEQ ID NO: 20
P11 ₍₃₁₉₋₃₃₃₎	HisAlaAsnPheCysLeuGlyProCysProTyrIleTrpSerLeu		SEQ ID NO: 1
P12 ₍₃₂₂₋₃₃₅₎	PheCysLeuGlyProCysProTyrIleTrpSerLeuAspThr		SEQ ID NO: 2
P13 ₍₃₂₆₋₃₃₉₎	ProCysProTyrIleTrpSerLeuAspThrGlnTyrSerLys		SEQ ID NO: 21
P14 ₍₃₃₀₋₃₄₃₎	IleTrpSerLeuAspThrGlnTyrSerLysValLeuAlaLeu		SEQ ID NO: 22
P15 ₍₃₃₅₋₃₄₉₎	ThrGlnTyrSerLysValLeuAlaLeuTyrAsnGlnHisAsnPro		SEQ ID NO: 23
P16 ₍₃₃₆₋₃₄₉₎	GlnTyrSerLysValLeuAlaLeuTyrAsnGlnHisAsnPro		SEQ ID NO: 24
P17 ₍₃₄₀₋₃₅₃₎	ValLeuAlaLeuTyrAsnGlnHisAsnProGlyAlaSerAla		SEQ ID NO: 25
P18 ₍₃₄₃₋₃₅₈₎	LeuTyrAsnGlnHisAsnProGlyAlaSerAlaAlaProCysCys		SEQ ID NO: 26
P19 ₍₃₄₄₋₃₅₈₎	TyrAsnGlnHisAsnProGlyAlaSerAlaAlaProCysCys		SEQ ID NO: 27
P20 ₍₃₄₈₋₃₆₀₎	AsnProGlyAlaSerAlaAlaProCysCysValProGln		SEQ ID NO: 28
P21 ₍₃₅₀₋₃₆₃₎	GlyAlaSerAlaAlaProCysCysValProGlnAlaLeuGlu		SEQ ID NO: 29
P22 ₍₃₅₄₋₃₆₇₎	AlaProCysCysValProGlnAlaLeuGluProLeuProIle		SEQ ID NO: 30
P23 ₍₃₅₈₋₃₇₁₎	ValProGlnAlaLeuGluProLeuProIleValTyrTyrVal		SEQ ID NO: 31
P24 ₍₃₆₄₋₃₇₇₎	ProLeuProIleValTyrTyrValGlyArgLysProLysVal		SEQ ID NO: 32
P25 ₍₃₆₈₋₃₈₁₎	ValTyrTyrValGlyArgLysProLysValGluGlnLeuSer		SEQ ID NO: 33
P26 ₍₃₇₂₋₃₈₅₎	GlyArgLysProLysValGluGlnLeuSerAsnMetIleVal		SEQ ID NO: 34
P27 ₍₃₇₈₋₃₉₁₎	GluGlnLeuSerAsnMetIleValArgSerCysLysCysSer		SEQ ID NO: 35

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Page 25, first paragraph, rewrite as follows:

C2

<u>Peptide</u>	<u>Sequence</u>	
P12 ₍₃₂₂₋₃₃₃₎	PheCysLeuGlyProCysProTyrIleTrpSerLeuAspThr	SEQ ID NO: 2
P28 ₍₃₂₂₋₃₄₄₎	PheCysLeuGlyProCysProTyrIleTrpSerLeuAspThrGlnLysVal LeuAlaLeuTyr	SEQ ID NO: 36
P29 ₍₃₁₃₋₃₅₅₎	HisGluProLysGlyTyrHisAlaAsnPheCysLeuGlyProCysProTyr IleTrpSerLeuAspThr	SEQ ID NO: 10
P30	PheSerLeuGlyProCysProTyrIleTrpSerLeuAspThr	SEQ ID NO: 37
P31	PheCysLeuGlyProSerProTyrIleTrpSerLeuAspThr	SEQ ID NO: 38
P32	PheSerLeuGlyProSerProTyrIleTrpSerLeuAspThr	SEQ ID NO: 39
P33	PheCysLeuGlyProCysProTyrIleTrpSerAspAspAsp	SEQ ID NO: 40
P34	AspAspAspGlyProCysProTyrIleTrpSerLeuAspThr	SEQ ID NO: 41
P35	AspAspAspGlyProCysProTyrIleTrpSerAspAspAsp	SEQ ID NO: 42
P36	GlyProCysProTyrIleTrpSerAspAspAsp	SEQ ID NO: 43
P37	AspAspAspGlyProCysProTyrIleTrpSer	SEQ ID NO: 44
P38	AspGlyProCysProTyrIleTrpSerAsp	SEQ ID NO: 45

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Paragraph beginning on page 28 on line 1 and ending on page 30, last line, rewrite as follows:

<u>Peptide</u>	<u>Sequence</u>	
P39 ₍₁₉₁₋₁₀₂₎	AsnProIleAlaSerValHisThrHisHisLysPro	SEQ ID NO: 46
P40 ₍₁₀₄₋₁₁₅₎	ValPheLeuLeuAsnSerProGlnProLeuValTrp	SEQ ID NO: 47
P41 ₍₁₀₉₋₁₂₀₎	SerProGlnProLeuValTrpHisLeuLysThrGlu	SEQ ID NO: 48
P42 ₍₁₁₀₋₁₂₁₎	ProGlnProLeuValTrpHisLeuLysThrGluArg	SEQ ID NO: 49
P43 ₍₃₃₃₋₃₄₄₎	TrpAlaLeuAspAsnGlyTyrArgProValThrSer	SEQ ID NO: 50
P44 ₍₄₂₈₋₄₃₉₎	ProIleValProSerValGlnLeuLeuProAspHis	SEQ ID NO: 51
P45 ₍₅₅₅₋₅₆₆₎	GlyAspGluGlyGluThrAlaProLeuSerArgAla	SEQ ID NO: 52
P46 ₍₅₆₃₋₅₇₄₎	LeuSerArgAlaGlyValValPheAsnCysSer	SEQ ID NO: 53
P47 ₍₆₀₃₋₆₁₄₎	LeuPheLeuValProSerProGlyValPheSerVal	SEQ ID NO: 54
P48 ₍₆₀₅₋₆₁₆₎	LeuValProSerProGlyValPheSerValAlaGlu	SEQ ID NO: 55
P49 ₍₇₀₇₋₇₁₈₎	GluLeuThrLeuCysSerArgLysLysGlySerLeu	SEQ ID NO: 56
P50 ₍₇₁₂₋₇₂₃₎	SerArgLysLysGlySerLeuLysLeuProArgCys	SEQ ID NO: 57
P51 ₍₇₁₇₋₇₂₈₎	SerLeuLysLeuProArgCysValThrProAspAsp	SEQ ID NO: 58
P52 ₍₇₂₂₋₇₃₃₎	ArgCysValThrProAspAspAlaCysThrSerLeu	SEQ ID NO: 59
P53 ₍₇₂₇₋₇₃₈₎	AspAspAlaCysThrSerLeuAspAlaThrMetIle	SEQ ID NO: 60
P54 ₍₇₃₁₋₇₄₂₎	ThrSerLeuAspAlaThrMetIleTrpThrMetMet	SEQ ID NO: 3
P55 ₍₇₃₂₋₇₄₃₎	SerLeuAspAlaThrMetIleTrpThrMetMetGln	SEQ ID NO: 61
P56 ₍₇₃₇₋₇₄₈₎	MetIleTrpThrMetMetGlnAsnLysLysThrPhe	SEQ ID NO: 62
P57 ₍₇₄₂₋₇₅₂₎	MetGlnAsnLysLysThrPheThrLysProLeuAla	SEQ ID NO: 63
P58 ₍₇₄₇₋₇₅₈₎	ThrPheThrLysProLeuAlaValValLeuGlnVal	SEQ ID NO: 64
P59 ₍₇₆₁₋₇₇₅₎	LysGluAsnValProSerThrLysAspSerSerProIleProPro	SEQ ID NO: 65
P60 ₍₇₆₆₋₇₈₀₎	SerThrLysAspSerSerProIleProProProProGlnIle	SEQ ID NO: 66
P61 ₍₇₇₁₋₇₈₅₎	SerProIleProProProProGlnIlePheHisGlyLeuAsp	SEQ ID NO: 67
P62 ₍₇₇₆₋₇₉₀₎	ProProProGlnIlePheHisGlyLeuAspThrLeuThrValMet	SEQ ID NO: 68
P63 ₍₇₈₁₋₇₉₅₎	PheHisGlyLeuAspThrLeuThrValMetGlyIleAlaPheAla	SEQ ID NO: 69
P64 ₍₇₈₆₋₈₀₀₎	ThrLeuThrValMetGlyIleAlaPheAlaAlaPheValIleGly	SEQ ID NO: 70
P65 ₍₇₉₇₋₈₀₉₎	LeuLeuThrGlyAlaLeuTrpTyrIleTyrSerHis	SEQ ID NO: 71
P66 ₍₄₅₋₅₉₎	LeuMetGluSerPheThrValLeuSerGlyCysAlaSerArgGly	SEQ ID NO: 72
P67 ₍₅₀₋₆₄₎	ThrValLeuSerGlyCysAlaSerArgGlyThrThrGlyLeuPro	SEQ ID NO: 73
P68 ₍₅₅₋₆₉₎	CysAlaSerArgGlyThrThrGlyLeuProArgGluValHisVal	SEQ ID NO: 74
P69 ₍₆₀₋₇₄₎	ThrThrGlyLeuProArgGluValHisValLeuAsnLeuArgSer	SEQ ID NO: 75
P70 ₍₆₅₋₇₉₎	ArgGluValHisValLeuAsnLeuArgSerThrAspGlnGlyPro	SEQ ID NO: 76
P71 ₍₇₀₋₈₄₎	LeuAsnL uArgSerThrAspGlnGlyProGlyGlnArgGlnArg	SEQ ID NO: 77
P72 ₍₇₅₋₈₉₎	ThrAspGlnGlyProGlyGlnArgGlnArgGluValThrLeuHis	SEQ ID NO: 78
P73 ₍₈₀₋₉₄₎	GlyGlnArgGlnArgGluValThrLeuHisLeuAsnProIleAla	SEQ ID NO: 79

P74 ₍₈₅₋₉₉₎	GluValThrLeuHisLeuAsnProIleAlaSerValHisThrHis	SEQ ID NO: 80
P75 ₍₉₀₋₁₀₄₎	LeuAsnProIleAlaSerValHisThrHisHisLysProIleVal	SEQ ID NO: 81
P76 ₍₉₅₋₁₀₉₎	SerValHisThrHisHisLysProIleValPheLeuLeuAsnSer	SEQ ID NO: 82
P77 ₍₁₀₀₋₁₁₄₎	HisLysProIleValPheLeuLeuAsnSerProGlnProLeuVal	SEQ ID NO: 83
P78 ₍₁₀₅₋₁₁₉₎	PheLeuLeuAsnSerProGlnProLeuValTrpHisLeuLysThr	SEQ ID NO: 84
P79 ₍₁₁₀₋₁₂₄₎	ProGlnProLeuValTrpHisLeuLysThrGluArgLeuAlaAla	SEQ ID NO: 85
P80 ₍₁₁₅₋₁₂₉₎	TrpHisLeuLysThrGluArgLeuAlaAlaGlyValProArgLeu	SEQ ID NO: 86
P81 ₍₁₂₀₋₁₃₄₎	ArgLeuAlaAlaGlyValProArgLeuPheLeuValSerGluGly	SEQ ID NO: 87
P82 ₍₁₂₅₋₁₃₉₎	GlyValProArgLeuPheLeuValSerGluGlySerValValGln	SEQ ID NO: 88
P83 ₍₁₃₀₋₁₄₄₎	PheLeuValSerGluGlySerValValGlnPheProSerGlyAsn	SEQ ID NO: 89
P84 ₍₁₃₅₋₁₄₉₎	GlySerValValGlnPheProSerGlyAsnPheSerLeuThrAla	SEQ ID NO: 90
P85 ₍₁₄₀₋₁₅₄₎	PheProSerGlyAsnPheSerLeuThrAlaGluThrGluGluArg	SEQ ID NO: 91
P86 ₍₁₄₅₋₁₅₉₎	PheSerLeuThrAlaGluThrGluGluArgAsnPheProGlnGlu	SEQ ID NO: 92
P87 ₍₁₅₀₋₁₆₄₎	GluThrGluGluArgAsnPheProGlnGluAsnGluHisLeuVal	SEQ ID NO: 93
P88 ₍₁₅₅₋₁₆₉₎	AsnPheProGlnGluAsnGluHisLeuValArgTrpAlaGlnLys	SEQ ID NO: 94
P89 ₍₁₆₀₋₁₇₄₎	AsnGluHisLeuValArgTrpAlaGlnLysGluTyrGlyAlaVal	SEQ ID NO: 95
P90 ₍₁₆₅₋₁₇₉₎	ArgTrpAlaGlnLysGluTyrGlyAlaValThrSerPheThrGlu	SEQ ID NO: 96
P91 ₍₁₇₀₋₁₈₄₎	GluTyrGlyAlaValThrSerPheThrGluLeuLysIleAlaArg	SEQ ID NO: 97
P92 ₍₁₇₅₋₁₈₉₎	ThrSerPheThrGluLeuLysIleAlaArgAsnIleTyrIleLys	SEQ ID NO: 98
P93 ₍₁₈₀₋₁₉₄₎	LeuLysIleAlaArgAsnIleTyrIleLysValGlyGluAspGln	SEQ ID NO: 99
P94 ₍₁₈₅₋₁₉₉₎	AsnIleTyrIleLysValGlyGluAspGlnValPheProProThr	SEQ ID NO: 100
P95 ₍₁₉₀₋₂₀₄₎	ValGlyGluAspGlnValPheProProThrCysAsnIleGlyLys	SEQ ID NO: 101
P96 ₍₁₉₅₋₂₀₉₎	ValPheProProThrCysAsnIleGlyLysAsnPheLeuSerLeu	SEQ ID NO: 102
P97 ₍₂₀₀₋₂₁₄₎	CysAsnIleGlyLysAsnPheLeuSerLeuAsnTyrLeuAlaGlu	SEQ ID NO: 103
P98 ₍₂₀₅₋₂₁₉₎	AsnPheLeuSerLeuAsnTyrLeuAlaGluTyrLeuGlnProLys	SEQ ID NO: 104
P99 ₍₂₁₀₋₂₂₄₎	AsnTyrLeuAlaGluTyrLeuGlnProLysAlaAlaGluGlyCys	SEQ ID NO: 105
P100 ₍₂₁₅₋₂₂₉₎	TyrLeuGlnProLysAlaAlaGluGlyCysValLeuProSerGln	SEQ ID NO: 106
P101 ₍₂₂₀₋₂₃₄₎	AlaAlaGluGlyCysValLeuProSerGlnProHisGluLysGlu	SEQ ID NO: 107
P102 ₍₂₂₅₋₂₃₉₎	ValLeuProSerGlnProHisGluLysGluValHisIleIleGlu	SEQ ID NO: 108
P103 ₍₂₃₀₋₂₄₄₎	ProHisGluLysGluValHisIleIleGluLeuIleThrProSer	SEQ ID NO: 109
P104 ₍₂₃₅₋₂₄₉₎	ValHisIleIleGluLeuIleThrProSerSerAsnProTyrSer	SEQ ID NO: 110
P105 ₍₂₄₀₋₂₅₄₎	LeuIleThrProSerSerAsnProTyrSerAlaPheGlnValAsp	SEQ ID NO: 111
P110 ₍₂₆₅₋₂₇₉₎	AspProGluValValLysAsnLeuValLeuIleLeuLysCysLys	SEQ ID NO: 115
P111 ₍₂₇₀₋₂₈₄₎	LysAsnLeuValLeuIleLeuLysCysLysLysSerValAsnTrp	SEQ ID NO: 116
P112 ₍₂₇₅₋₂₈₉₎	IleLeuLysCysLysLysSerValAsnTrpValIleLysSerPhe	SEQ ID NO: 117
P113 ₍₂₈₀₋₂₉₄₎	LysSerValAsnTrpValIleLysSerPheAspValLysGlyAsn	SEQ ID NO: 118
P114 ₍₂₈₅₋₂₉₉₎	ValIleLysSerPheAspValLysGlyAsnLeuLysValIleAla	SEQ ID NO: 119
P115 ₍₂₉₀₋₃₀₄₎	AspValLysGlyAsnLeuLysValIleAlaProAsnSerIleGly	SEQ ID NO: 120

P106 ₍₂₄₅₋₂₅₉₎	SerAsnProTyrSerAlaPheGlnValAspIleIleValAspIle	SEQ ID NO: 4
P107 ₍₂₅₀₋₂₆₄₎	AlaPheGlnValAspIleIleValAspIleArgProAlaGlnGlu	SEQ ID NO: 112
P108 ₍₂₅₅₋₂₆₉₎	IleIleValAspIleArgProAlaGlnGluAspProGluValVal	SEQ ID NO: 113
P109 ₍₂₆₀₋₂₇₄₎	ArgProAlaGlnGluAspProGluValValLysAsnLeuValLeu	SEQ ID NO: 114
P116 ₍₂₉₅₋₃₀₉₎	LeuLysValIleAlaProAsnSerIleGlyPheGlyLysGluSer	SEQ ID NO: 121
P117 ₍₃₀₀₋₃₁₄₎	ProAsnSerIleGlyPheGlyLysGluSerGluArgSerMetThr	SEQ ID NO: 122
P118 ₍₃₀₅₋₃₁₉₎	PheGlyLysGluSerGluArgSerMetThrMetThrLysLeuVal	SEQ ID NO: 123
P119 ₍₃₁₀₋₃₂₄₎	GluArgSerMetThrMetThrLysLeuValArgAspAspIlePro	SEQ ID NO: 124
P120 ₍₃₁₅₋₃₂₉₎	MetThrLysLeuValArgAspAspIleProSerThrGlnGluAsn	SEQ ID NO: 125
P121 ₍₃₂₀₋₃₃₄₎	ArgAspAspIleProSerThrGlnGluAsnLeuMetLysTrpAla	SEQ ID NO: 126
P122 ₍₃₂₅₋₃₃₉₎	SerThrGlnGluAsnLeuMetLysTrpAlaLeuAspAsnGlyTyr	SEQ ID NO: 127
P123 ₍₃₃₀₋₃₄₄₎	LeuMetLysTrpAlaLeuAspAsnGlyTyrArgProValThrSer	SEQ ID NO: 128
P124 ₍₃₃₅₋₃₄₉₎	LeuAspAsnGlyTyrArgProValThrSerTyrThrMetAlaPro	SEQ ID NO: 129
P125 ₍₃₄₀₋₃₅₄₎	ArgProValThrSerTyrThrMetAlaProValAlaAsnArgPhe	SEQ ID NO: 130
P126 ₍₃₄₅₋₃₅₉₎	TyrThrMetAlaProValAlaAsnArgPheHisLeuArgLeuGlu	SEQ ID NO: 131
P127 ₍₃₅₀₋₃₆₄₎	ValAlaAsnArgPheHisLeuArgLeuGluAsnAsnGluGluMet	SEQ ID NO: 132
P128 ₍₃₅₅₋₃₆₉₎	HisLeuArgLeuGluAsnAsnGluGluMetArgAspGluGluVal	SEQ ID NO: 133
P129 ₍₃₆₀₋₃₇₄₎	AsnAsnGluGluMetArgAspGluGluValHisThrIleProPro	SEQ ID NO: 134
P130 ₍₃₆₅₋₃₇₉₎	ArgAspGluGluValHisThrIleProProGluLeuArgIleLeu	SEQ ID NO: 135
P131 ₍₃₇₀₋₃₈₄₎	HisThrIleProProGluLeuArgIleLeuLeuAspProAspHis	SEQ ID NO: 136
P132 ₍₃₇₅₋₃₈₉₎	GluLeuArgIleLeuLeuAspProAspHisProProAlaLeuAsp	SEQ ID NO: 137
P133 ₍₃₈₀₋₃₉₄₎	LeuAspProAspHisProProAlaLeuAspAsnProLeuPhePro	SEQ ID NO: 138
P134 ₍₃₈₅₋₃₉₉₎	ProProAlaLeuAspAsnProLeuPheProGlyGluGlySerPro	SEQ ID NO: 139
P135 ₍₃₉₀₋₄₀₄₎	AsnProLeuPheProGlyGluGlySerProAsnGlyGlyLeuPro	SEQ ID NO: 140
P136 ₍₃₉₅₋₄₀₉₎	GlyGluGlySerProAsnGlyGlyLeuProPheProPheProAsp	SEQ ID NO: 141
P137 ₍₄₀₀₋₄₁₄₎	AsnGlyGlyLeuProPheProPheProAspIleProArgArgGly	SEQ ID NO: 142
P138 ₍₄₀₅₋₄₁₉₎	PheProPheProAspIleProArgArgGlyTrpLysGluGlyGlu	SEQ ID NO: 143

C3
Contd.

Page 32, first full paragraph, rewrite as follows:

C4

Table 5. Peptides derived from modification of peptide P54 (peptides P139 to P143) and of the human type III receptor (peptides P144 and P145).

Peptide	Sequence	Derivation	
P54 ₍₇₃₁₋₇₄₂₎	ThrSerLeuAspAlaThrMetIleTrpThrMetMet	Rat type III receptor	SEQ ID NO: 3
P139	ThrSerLeuAspAlaThrMetIleTrpAspAspAsp		SEQ ID NO: 144
P140	AspAspAspAlaThrMetIleTrpThrMetMet		SEQ ID NO: 145
P141	AspAlaThrMetIleTrpAsp		SEQ ID NO: 146
P142	ThrSerLeuMetIleTrpThrMetMet		SEQ ID NO: 5
P143	ThrSerLeuAspAlaThrThrMetMet		SEQ ID NO: 147
P144 ₍₁₃₁₋₁₄₂₎	ThrSerLeuAspAlaSerIleIleTrpAlaMetMet GlnAsn	Human type III receptor	SEQ ID NO: 6
P145 ₍₂₄₁₋₂₅₄₎	SerAsnProTyrSerAlaPheGlnValAspIleThr IleAsp	Human type III receptor	SEQ ID NO: 7

Paragraph beginning on page 34, line 3 and ending on page 35, line 8, rewrite as follows:

C5

Peptide	Sequence	Origin	
P146 ₍₈₄₋₁₀₁₎	CysValAlaValTrpArgLysAsnAspGluAsnIleThr LeuGluThrValCys	Type II receptor	SEQ ID NO: 148
P147 ₍₁₁₄₋₁₂₂₎	CysAspPheGlnLeuLeuLysLeuAspGlyLysPheSer ValValTyrAlaLysCys	Fetuin	SEQ ID NO: 149
P148 ₍₁₁₄₋₁₂₂₎	CysAspPheHisIleLeuLysGlnAspGlyGlnPheArg ValCysHisAlaGlnCys	Fetuin	SEQ ID NO: 150
P149 ₍₁₁₄₋₁₂₂₎	CysAspIleHisValLeuLysGlnAspGlyPheSerVal LeuPheThrLysCysAsp	Fetuin	SEQ ID NO: 151
P150 ₍₂₄₇₋₂₆₁₎	GluAlaValLeuIleLeuGlnGlyProProTyrValSer TrpLeu	Endoglin	SEQ ID NO: 8
P151 ₍₃₈₉₋₃₉₃₎	ValAsnLeuProAspThrArgGlnGlyLeuLeuGluGlu AlaArg	Endoglin	SEQ ID NO: 152
P152 ₍₄₄₅₋₄₅₅₎	LeuAspSerLeuSerPheGlnLeuGlyLeuTyrLeuSer ProHis	Endoglin	SEQ ID NO: 9
P153 ₍₄₆₁₋₄₅₅₎	ProSerIleProGluLeuMetThrGlnLeuAspSerCys GlnLeu	Endoglin	SEQ ID NO: 153
P154 ₍₄₇₉₋₄₅₅₎	MetSerProSerIleProGluLeuMetThrGlnLeuAsp SerCys	Endoglin	SEQ ID NO: 154
P155 ₍₁₃₋₂₄₎	LeuLeuLeuValLeuLeuProThrAspAlaSer	α -2-Macroglobulin	SEQ ID NO: 155
P156 ₍₂₀₋₃₁₎	ProThrAspAlaSerValSerGlyLysProGlnTyr	α -2-Macroglobulin	SEQ ID NO: 156
P157 ₍₄₄₋₅₅₎	ThrGluLysGlyCysValLeuLeuSerTyrLeuAsn	α -2-Macroglobulin	SEQ ID NO: 157
P158 ₍₁₆₆₋₁₇₇₎	TyrIleGlnAspProLysGlyAsnArgIleAlaGln	α -2-Macroglobulin	SEQ ID NO: 158
P158 ₍₁₆₆₋₁₇₇₎	TyrIleGlnAspProLysGlyAsnArgIleAlaGln	α -2-Macroglobulin	SEQ ID NO: 158
P159 ₍₁₉₃₋₂₀₃₎	PheProLeuSerSerGluProPheGlnGlySerTyr	α -2-Macroglobulin	SEQ ID NO: 159
P160 ₍₂₄₇₋₂₅₈₎	AsnValSerValCysGlyLeuTyrThrTyrGlyLys	α -2-Macroglobulin	SEQ ID NO: 160
P161 ₍₂₄₆₋₂₅₉₎	ValSerValCysGlyLeuTyrThrTyrGlyLysPro	α -2-Macroglobulin	SEQ ID NO: 161
P162 ₍₂₅₆₋₂₆₂₎	ValCysGlyLeuTyrThrTyrGlyLysProValPro	α -2-Macroglobulin	SEQ ID NO: 162
P163 ₍₂₆₇₋₂₇₈₎	SerIleCysArgLysTyrSerAspAlaSerAspCys	α -2-Macroglobulin	SEQ ID NO: 163
P164 ₍₄₆₉₋₄₈₀₎	ProCysGlyHisThrGlnThrValGlnAlaHisTyr	α -2-Macroglobulin	SEQ ID NO: 164
P165 ₍₅₅₄₋₅₆₃₎	AspSerAlaLysTyrAspValGluAsnCysLeuAla	α -2-Macroglobulin	SEQ ID NO: 165
P167 ₍₇₉₀₋₈₀₂₎	GlnProPhePheValGluLeuThrMetProTyrSer	α -2-Macroglobulin	SEQ ID NO: 167
P168 ₍₈₂₇₋₈₃₈₎	GlnLeuGluAlaSerProAlaPheLeuAlaValPro	α -2-Macroglobulin	SEQ ID NO: 168
P169 ₍₈₃₅₋₈₃₈₎	SerValGlnLeuGluAlaSerProAlaPheLeuAla	α -2-Macroglobulin	SEQ ID NO: 169
P170 ₍₈₇₆₋₈₈₇₎	AlaLeuGluSerGlnGluLeuCysGlyThrGluVal	α -2-Macroglobulin	SEQ ID NO: 170
P171 ₍₁₀₀₁₋₁₀₁₂₎	LysSerLysIleGlyTyrLeuAsnThrGlyTyr	α -2-Macroglobulin	SEQ ID NO: 171

P172 ₍₁₂₀₁₅₋₁₂₀₁₆₎	IleGlyTyrLeuAsnThrGlyTyrGlnArgGlnLeu	α -2-Macroglobulin	SEQ ID NO: 172
P173 ₍₁₂₀₂₁₋₁₂₀₂₂₎	LysArgLysGluValLeuLysSerLeuAsnGluGlu	α -2-Macroglobulin	SEQ ID NO: 173
P174 ₍₁₂₀₂₃₋₁₂₀₂₄₎	ValGlyHisPheTyrGluProGlnAlaProSerAla	α -2-Macroglobulin	SEQ ID NO: 174
P175 ₍₁₂₀₂₉₋₁₂₀₃₀₎	ThrSerTyrValLeuLeuAlaTyrLeuThrGlnAla	α -2-Macroglobulin	SEQ ID NO: 175
P176 ₍₁₂₀₃₁₋₁₂₀₃₂₎	TyrValLeuLeuAlaTyrLeuThrAlaGlnProAla	α -2-Macroglobulin	SEQ ID NO: 176
P177 ₍₁₂₀₃₆₋₁₂₀₃₇₎	ValAlaLeuHisAlaLeuSerLysTyrGlyAlaAla	α -2-Macroglobulin	SEQ ID NO: 177
P178 ₍₁₂₀₃₈₋₁₂₀₄₃₎	TyrGlyArgAsnGlnGlyAsnThrTrpLeuThrAla	α -2-Macroglobulin	SEQ ID NO: 178
P179 ₍₁₂₀₃₉₋₁₂₀₄₅₎	ArgAsnGlnGlyAsnThrTrpLeuThrAlaPheVal	α -2-Macroglobulin	SEQ ID NO: 179

Page 36, first full paragraph, rewrite as follows:

C6

Table 7. Comparison of the inhibitory activity of TGF β 1, of some peptides, measured by bioassay of inhibition of growth of the MV-1-Lu¹ cells (peptide concentration 200 μ g/ml) with inhibition of the binding of TGF β 1 to its cell receptors measured using flow cytometry² (peptide concentration 420 n μ /ml).

Peptides	Bioassay	Cytometry	Sequence	
	(% inhibition) ¹	(% inhibition) ²		
P29	77,6	92,34	HisGluProLysGlyTyrHis AlaAsnPheCysLeuGlyPro CysProTyrIleTrpSerLeu AspThr	SEQ ID NO: 10
P11	40	86	HisAlaAsnPheCysLeuGly ProCysProTyrIleTrpSer Leu	SEQ ID NO: 1
P12	96	77	PheCysLeuGlyProCysPro TyrIleTrpSerLeuAspThr	SEQ ID NO: 2
P18	18,2	6,5	LeuTyrAsnGlnHisAsnPro GlyAlaSerAlaAlaProCys Cys	SEQ ID NO: 26
P54	97	82,3	ThrSerLeuAspAlaThrMet IleTrpThrMetMet	SEQ ID NO: 3
P140	-1,7	69,8	AspAspAspAlaThrMetIle TrpThrMetMet	SEQ ID NO: 145
P142	70	72	ThrSerLeuMetIleTrpThr MetMet	SEQ ID NO: 5
P106	40	91	SerAsnProTyrSerAlaPhe GlnValAspIleIleValAsp Ile	SEQ ID NO: 4
P145	21	74,35	SerAsnProTyrSerAlaPhe GlnValAspIleThrIleAsp	SEQ ID NO: 7
P144	88	80	ThrSerLeuAspAlaSerIle IleTrpAlaMetMetGlnAsn	SEQ ID NO: 6
P150	64	73	GluAlaValLeuIleLeuGln GlyProProTyrValSerTrp Leu	SEQ ID NO: 8
P152	45	68,4	LeuAspSerLeuSerPheGln LeuGlyLeuTyrLeuSerPro His	SEQ ID NO: 9